# Creating your own functions Learning the basics of R - Part 3

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# Outline

- Why write functions
- When to write functions
- How to write functions
- Practical session

# Why write functions

- Allow automation of common tasks in a more powerful and general way than *copy-and-pasting*;
  - You can give a function an evocative name that makes your code easier to understand;
  - As requirements change, you only need to update code in one place, instead of many.
  - You eliminate the chance of making incidental mistakes when you copy and paste (i.e. updating a variable name in one place, but not in another).
- Fewer global variables: When you run a function, the intermediate variables that it creates are not stored in your global environment. This saves memory and keeps your global environment cleaner.
- Better documentation: Well documented functions help the user understand the steps of your processing.
- Easier to maintain / edit: When you create a function for a repeated task, it is easy to edit that one function. Then every location in your code where that same task is performed is automatically updated.

## When to write functions

You should consider writing a function whenever you've copied and pasted a block of code more than twice (i.e. you now have three copies of the same code).

For example, take a look at this code.

What does it do?

```
df <- data.frame(
    a = c(1, 2, 1, 1, 1, 2, 1, 1, 2, 2),
    b = c(2, 2, 2, 1, 1, 1, 1, 1, 2, 1, 2),
    c = c(1, 2, 1, 1, 2, 1, 2, 1, 2, 2)
)

df$a <- ifelse(df$a == 2, 0, df$a)
df$b <- ifelse(df$b == 2, 0, df$b)
df$c <- ifelse(df$c == 2, 0, df$c)</pre>
```

### When to write functions

#### Original data:

#### Recoded data:

##		а	b	С	
##	1	1	0	1	
##	2	0	0	0	
##	3	1	0	1	
##	4	1	1	1	
##	5	1	1	0	
##	6	0	1	1	
##	7	1	1	0	
##	8	1	0	1	
##	9	0	1	0	
##	10	0	0	0	

This is a good example of when writing a function will be useful/beneficial.

### How to write functions

```
We can create a function called recode_values():
```

```
recode_values <- function(x) {
   ifelse(x == 2, 0, x)
}</pre>
```

And apply it to the same data as follows:

```
df$a <- recode_values(df$a)
df$b <- recode_values(df$b)
df$c <- recode_values(df$c)</pre>
```

We get:

```
##abc##1101##2000##3101##4111##5111##6011##7110##8101##9010##10000
```

### How to write functions

- You need to pick a name for the function. In the example I used recode\_values because this function recodes the values based on a specified rule (i.e., value of 2 is converted to 0).
- You list the inputs, or **arguments**, to the function inside function. Here we have just one argument. If we had more the call would look like function(x, y, z).
- You place the code you have developed in body of the function, a { block that immediately follows function(...).

### **Questions?**

### **Practical session**

We'll work through *Exercise 2 - Manipulating objects and creating new functions* in Practical R for Epidemiologists (https://practical-norg/exercise2.html) as a GitHub Classroom assignment

# Thank you!

Slides can be viewed at https://oxford-ihtm.io/open-reproducible-science/session4.html PDF version of slides can be downloaded at https://oxford-ihtm.io/open-reproducible-science/pdf/session4r-basics-part3.pdf

R scripts for slides available her